STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: *10/720, 843C*
Source: *IFW/b*
Date Processed by STIC: *4/12/06*

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY
FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT
MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO **REDUCE** ERRORED SEQUENCE LISTINGS, **PLEASE** USE THE **CHECKER VERSION 4.4.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:
http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (**<http://www.uspto.gov/ebc/efs/downloads/documents.htm>** , **EFS Submission User Manual** - ePAVE)
2. **U.S. Postal Service:** Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. **Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):**
   U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06

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        RAW SEQUENCE LISTING                    DATE: 04/12/2006
        PATENT APPLICATION:  US/10/720,843C      TIME: 10:49:22

        Input Set : E:\Solseqlst.txt
        Output Set: N:\CRF4\04122006\J720843C.raw

      3 <110> APPLICANT: Solulink
      4        Schwartz, David A.
      6 <120> TITLE OF INVENTION: HYDRAZINE-BASED AND CARBONYL-BASED
      7        BIFUNCTIONAL CROSSLINKING REAGENTS
     10 <130> FILE REFERENCE: SOL.003.DIV1
C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/720,843C
     13 <141> CURRENT FILING DATE: 2003-11-24
     15 <150> PRIOR APPLICATION NUMBER: 09/815,978
     16 <151> PRIOR FILING DATE: 2001-03-22
E--> 18 <160> NUMBER OF SEQ ID NOS: 60/191,186
E--> 19 <161> 2000-03-22
     21 <170> SOFTWARE: 1
E--> 23 <180> FastSEQ for Windows Version 4.0
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Does Not Comply Corrected Diskette Needed

ERRORED SEQUENCES

SEQUENCE LISTING

<110> Solulink
      Schwartz, David A.

<120> HYDRAZINE-BASED AND CARBONYL-BASED
      BIFUNCTIONAL CROSSLINKING REAGENTS

<130> SOL.003.DIV1

<140> 10/702,843 *10/720,843*
<141> 2003-11-24

charge to

<150> 09/815,978
<151> 2001-03-22

<1507 <160> 60/191,186
<1577 <161> 2000-03-22

<1607 <170> 1

<1707 <180> FastSEQ for Windows Version 4.0

<210> 1
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<212> DNA
<213> Artificial Sequence

<220>
<223> 25-mer phosphodiester oligonucleotide modified to
      incorporate a C6-aminolinker

<221> modified_base
<222> 1
<223> n= n-hexylamino thymine

<400> 1
nttttttagc ctaactgatg ccatg   25

Suggestion: Consult Sequence Rules for valid format

<1507 and <1577 are used for all prior application.

<1607 is used for total number of sequences   *data*

<1707 is used for software   *<1807 is invalid*

3

Input Set : **E:\Solseqlst.txt**
Output Set: **N:\CRF4\04122006\J720843C.raw**

L:12 M:270 C: Current Application Number differs, Replaced Current Application Number
L:19 M:250 E: Invalid Numeric Identifier, INVALID IDENTIFIER
L:23 M:250 E: Invalid Numeric Identifier, INVALID IDENTIFIER
L:34 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:38 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:39 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
L:18 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (60) Counted (1)